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Hosted By:  
U.S. Department of Health and Human Services  
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# Confined Placental Mosaicism In Infants with Fetal Growth Restriction

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# Confined Placental Mosaicism - Historical Perspective

- occurs in 1-2 % of first trimester CVS samples
- all chromosomal mosaicism in placental samples is not confined
- 1/3 represents true mosaicism

# Etiologies

- Post fertilization event confined to one cell line
- “Rescue” to diploidy of an originally trisomic conception

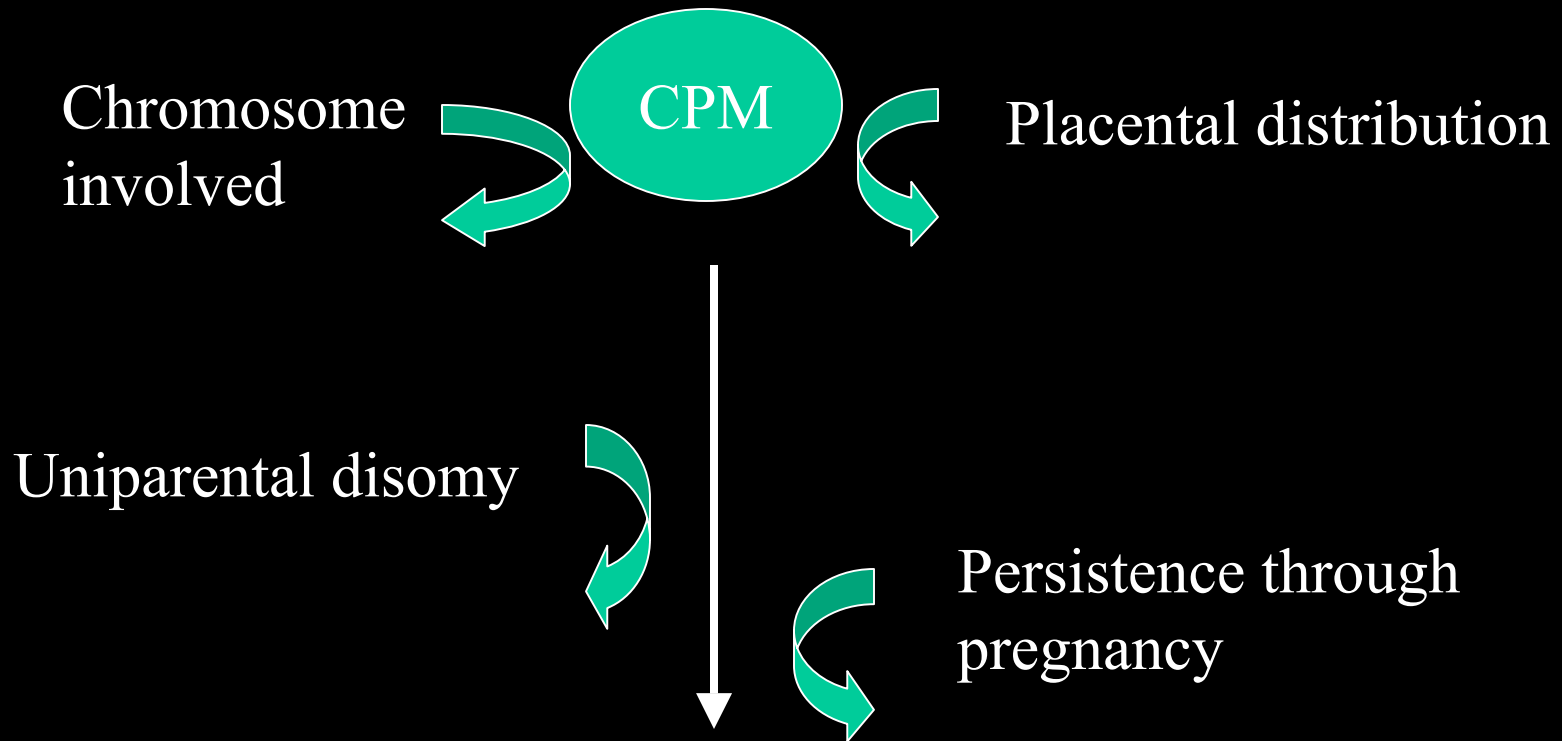
# Approaches to the study of CPM and fetal growth restriction

- Cohorts with CPM identified first trimester (CVS)
- Cohorts of newborns
- Case control studies of newborns

# Approach : follow-up of cohorts with CPM diagnosed first trimester

- Adverse outcomes suggested over 10 years ago – pregnancy loss, stillbirth, growth restriction

# Variation in outcomes



**FETAL GROWTH RESTRICTION**

# Variation in outcomes

## Chromosome involved

# 2,3,7,8 normal outcomes

# 9, 16, 22 of meiotic  
origin associated with  
IUGR

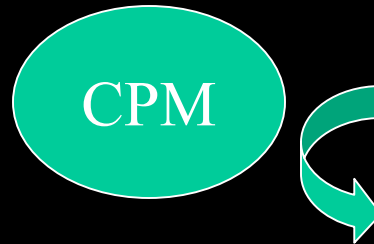
(Robinson, 1997)



**FETAL GROWTH RESTRICTION**



# Variation in outcomes



## Placental distribution

(Kalousek, 92; Simoni, 1994)

Type I – cytotrophoblast  
direct preparation

Type II – extraembryonic mesoderm  
culture preparation

Type III – both cell lines



**FETAL GROWTH RESTRICTION**

# Variation in outcomes



**Uniparental disomy**  
(Robinson, 1997)

<u>CPM16</u>	<u>IUGR</u>	<u>normal</u>
Fetal UPD	11	2
Fetal BPD	5	8

**FETAL GROWTH RESTRICTION**

# Variation in outcomes



**Persistence through pregnancy**

-variable 50-80%

-35% rate of IUGR  
(Kalousek, 1991)



**FETAL GROWTH RESTRICTION**

# Approach : CPM among a cohort of newborns (Artan, 1995)

- Karyotypes from 125 term placentas of pregnancies delivered following prenatal determination of normal fetal karyotype (AMA indication)
  - Higher risk population for nondisjunction
  - 6/125 (4.8%) CPM

- All 6 cases of CPM ended in IUGR infants
 

– 46,XX/47,XX,+14 (125/25)	2414	39 wks
– 46,XX/92,XXXX (74/76)	1647	34 wks
– 46,XY/47,XY, +21 (124/26)	2100	36 wks
– 46,XX/47,XX,+21 (73/87)	2400	40 wk
– 46,XX/45,X (61/79)	1760	38 wk
– 46,XY/47,XY,+18 (61/79)	2200	39 wks
- Birthweights CPM=2086+/-131.5;
  - normal placental biopsies 3305.2+/- 28.8

# Approach : analysis of growth restricted newborns - Unanswered Questions

- How large of a contributor is CPM to the population of infants with growth restriction?
- Are there characteristic clinical findings?

# Study Proposal for Case/control analysis— Primary Aim

- Determine the frequency of CPM by karyotype analysis of placental biopsies from infants with growth restriction compared to biopsies from placentas of maternal age matched, appropriately grown infants

# Study Proposal – Secondary Aims

- Utilize molecular, chromosome specific polymorphisms to identify uniparental disomy or low level mosaicism in a subset of patients if CPM not identified cytogenetically
- Explore clinical variables for identifying characteristics



# Background

- Which IUGR populations have been studied?
- Which chromosomes? Tetraploidy ?
- Alternative ways to search
  - Traditional cytogenetics
  - Molecular cytogenetics (FISH)
  - Molecular genetics (dinucleotide repeats)

# Studies of infants with unexplained IUGR

• Kalousek, 1983	2/9	
• Verp, 1990	0/11	
• Krishnamoorthy, 1995	4/26	
• Wilkins-Haug, 1995	3/12	
• Cowles, 1996	1/20	
• Stipolijev, 2001	<u>3/20</u>	
	13 / 98	(13.2 %)

# CPM among different populations of IUGR infants

Kennerknecht, 1993

- Newborns presenting with SGA 0/71
- Newborns having normal CVS  
who developed SGA (24/1300) 5/24
- Controls 0/20

# What do these studies suggest?

- CPM may play a role in the significantly IUGR population – those characterized by antepartum diagnoses, nonreassuring fetal well-being
- Sample sizes of both case and controls need to be adequate
- Role of tetraploidy ?

# Aneuploidy versus tetraploidy – Is there any evidence to support tetraploidy as a pathologic factor?

- Considered artifact - time in culture  
Tegenkamp, 1976; Kaji. 1979, 1981)

# Does tetraploidy occur “in vivo”?

- preimplantation embryos
- uncultured amnion by sex chromatin and cellular DNA determinations (Klinger, 1960)
- Tetraploidy by flow cytometry in placenta
  - 2.2% tetraploid

# Background rate of tetraploidy

(Noomen, 2001)

- 100 women AMA
- Semi direct and long term culture of chorionic villi
- Up to three tetraploids in 27% of STC
- In all long term cultures

# Any association of tetraploidy with abnormal placentation?

- Miscarriages assessed by long term culture (Hunt, 1985)
  - 10-30% in spontaneous miscarriages
  - 10% tetraploidy in first trimester tabs
- Miscarriages assessed by direct preparation (Eiben, 1990)
  - 9.2% tetraploidy



# Tetraploidy among CPM

- 5% of CPM is tetraploid mosaic (Ledbetter, 1992)
- ACC UK collaborative data (1994)
  - Tetraploidy noted as well

# Materials and Methods

- Antepartum identification of IUGR by ultrasound as  $<10\%$  for gestational age
- Singleton pregnancies with EDC confirmed by US  $< 16$  weeks gestation
- Excluded maternal conditions of HTN, IDDM, SLE, fetal malformations

# Sample Sizes

- 75 IUGR cases without recognized risk factors
- 75 AGA controls matched by maternal age to within 5 years
- 95% confidence with 80% power to detect  $\geq$  15% CPM among IUGR population
- Assumes 0.5 % CPM among AGA controls

# Study samples

- placental biopsies
- cord blood for karyotype or ability to recontact
- parental buccal samples or peripheral blood sample for DNA extraction

# Placental Samples

- paired chorionic plate samples removed from a mapped 4 locations
- one for culture
- one for disaggregated nuclei (FISH or DNA extaction)

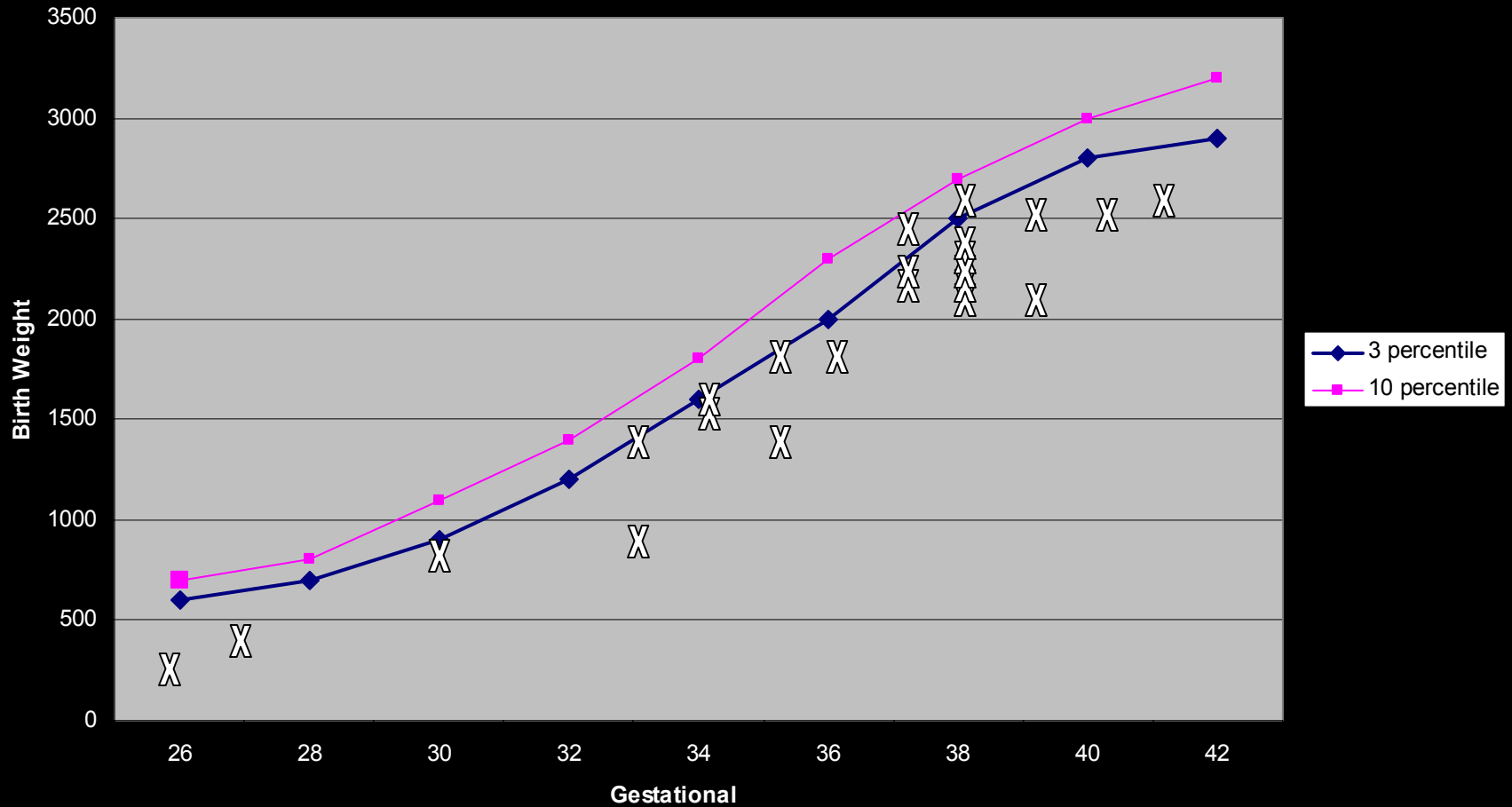
# Karyotype analysis

- Cultures established according to routine long term protocols
- 25 cells scored from each site (excludes > 15% mosaicism with 95% confidence)

# Molecular analysis

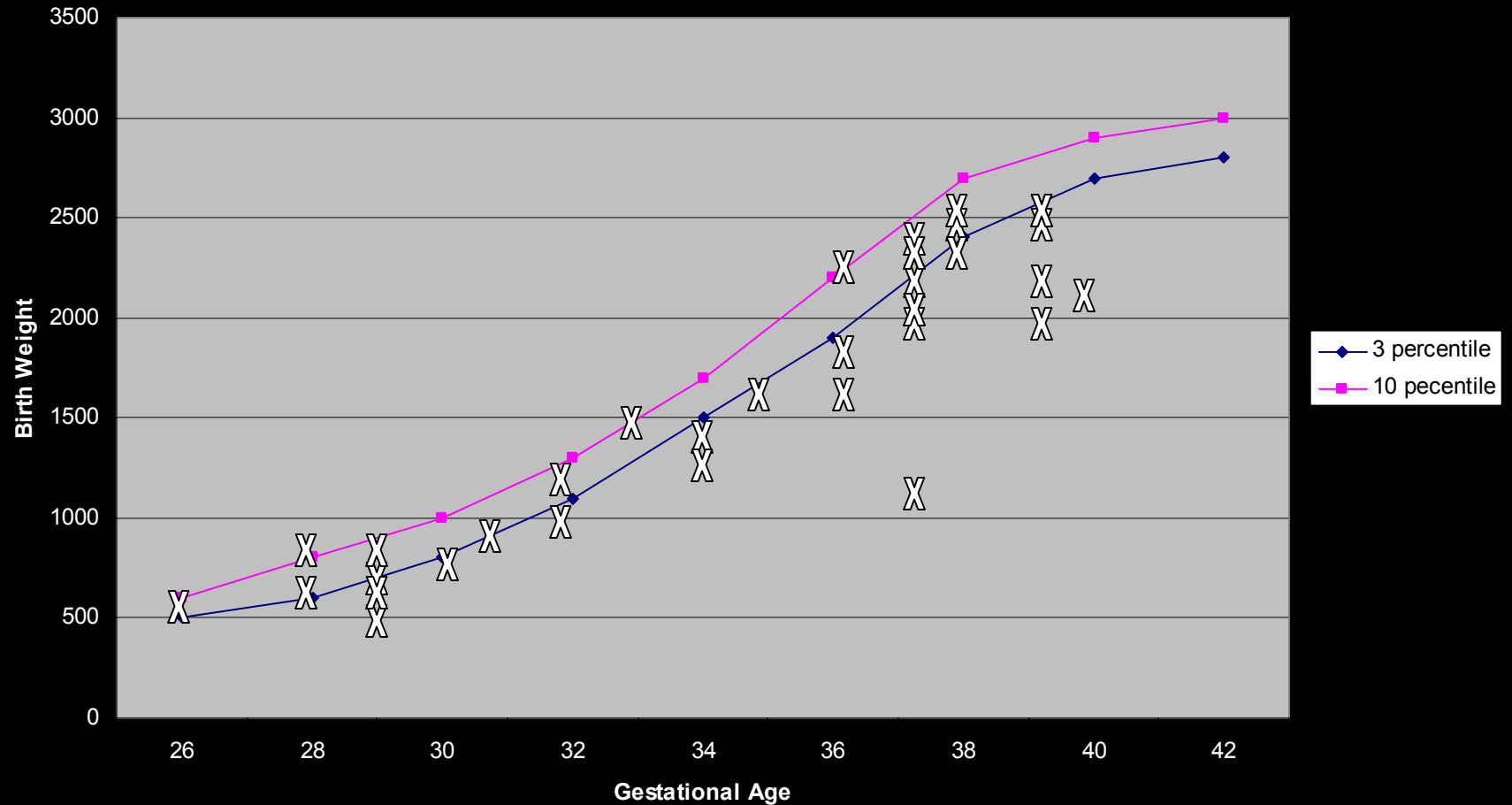
- Fluorescent panel of dinucleotide markers with heterozygosity scores of  $> 0.75$
- Automated genotyping on ABI377
- Minimum of 1 and maximum of three markers per each autosome

# Birth Weight Distribution - Males





# Birth Weight Distribution - Females



# Results

	<u>Aneuploid</u>	<u>Tetraploid</u>	<u>Total</u>
• Cases	1	5	6/75
• Controls	1	0	1/75

# Aneuploidy Mosaicism

- Case diploid/aneuploid  
– 46,XX/48,XX,+17,+21 14 / 12
- Control  
– 46,XY/47,XY,+10 7 / 13

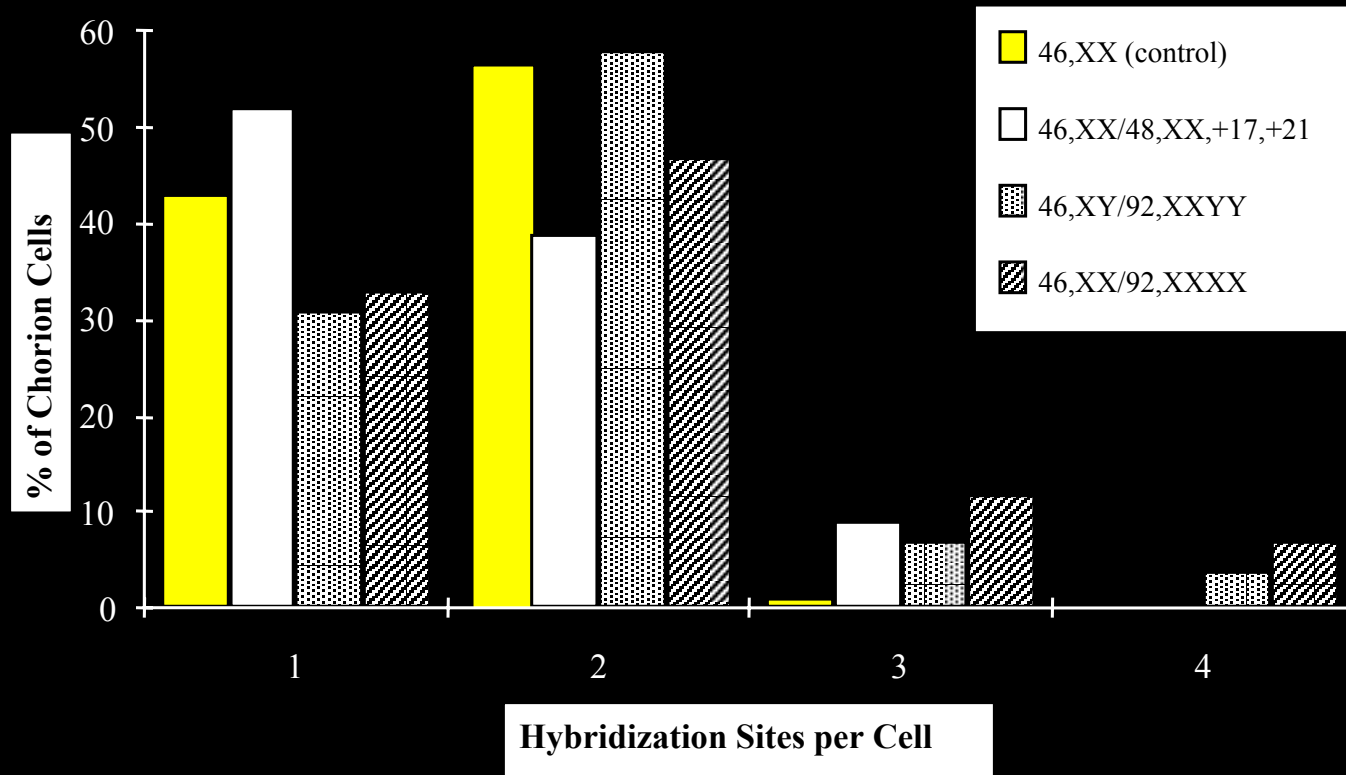
# Tetraploid Mosaicism

- Cases

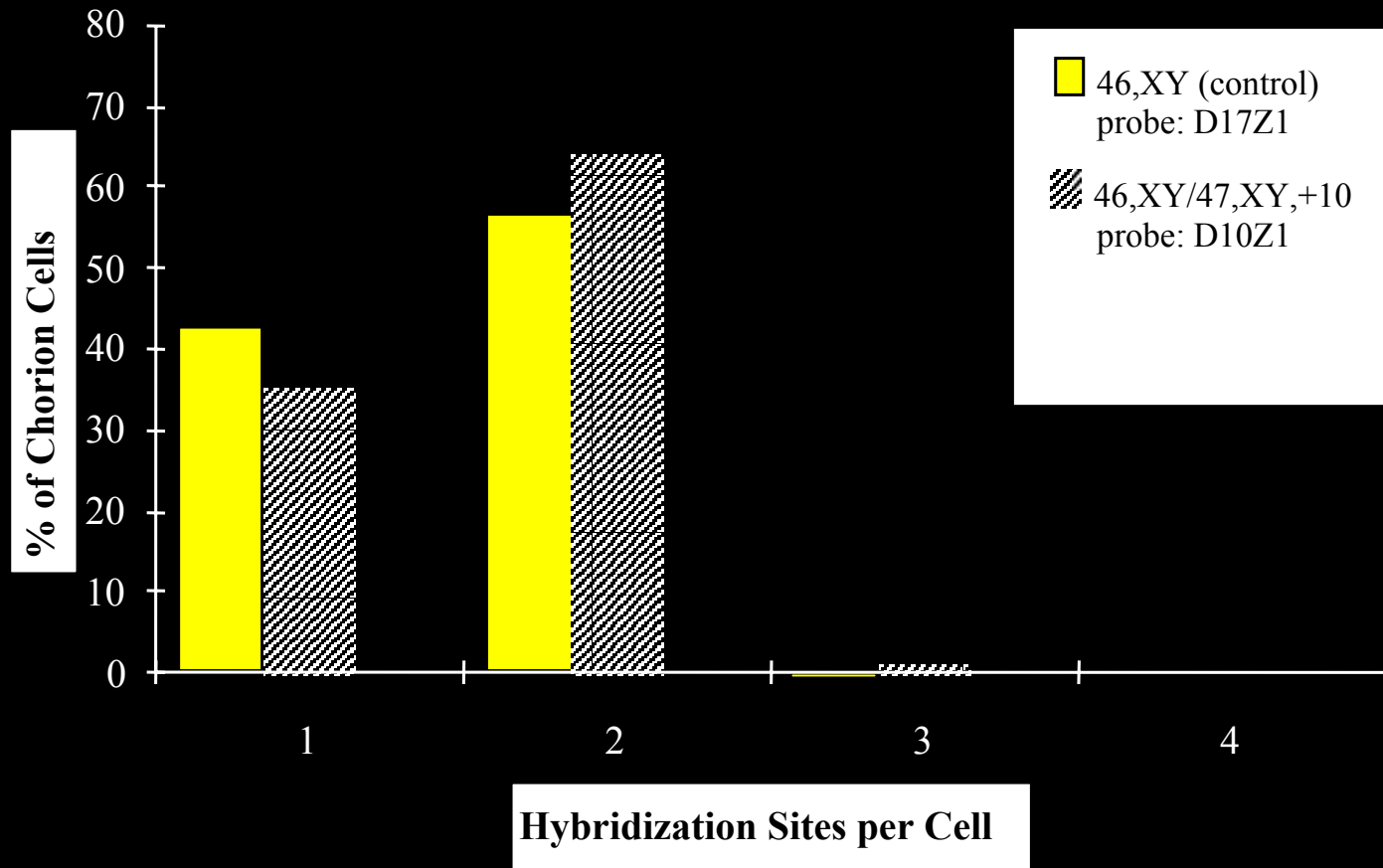
	diploid/polyploid	days in culture
– 46,XX/92,XXXX	25 / 30	10
– 46,XY/92,XXYY	25 / 13	13
– 46,XX/92,XXXX	25 / 23	10
– 46,XX/92,XXXX	25 / 10	8
– 46,XX/92,XXXX	25 / 19	12
- Controls
  - none

# Hybridization Sites in IUGR Placentas

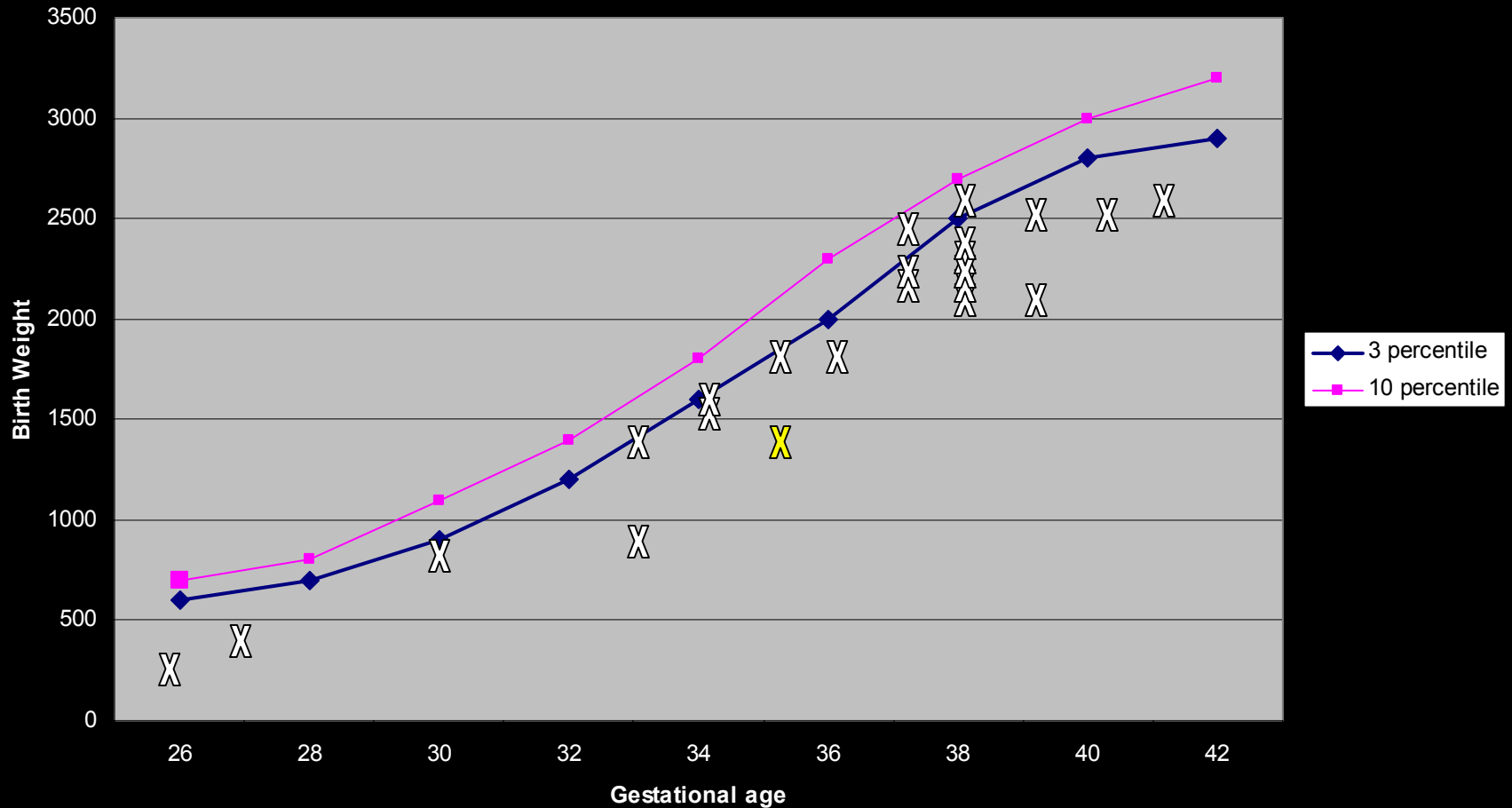
Probe:D17Z1



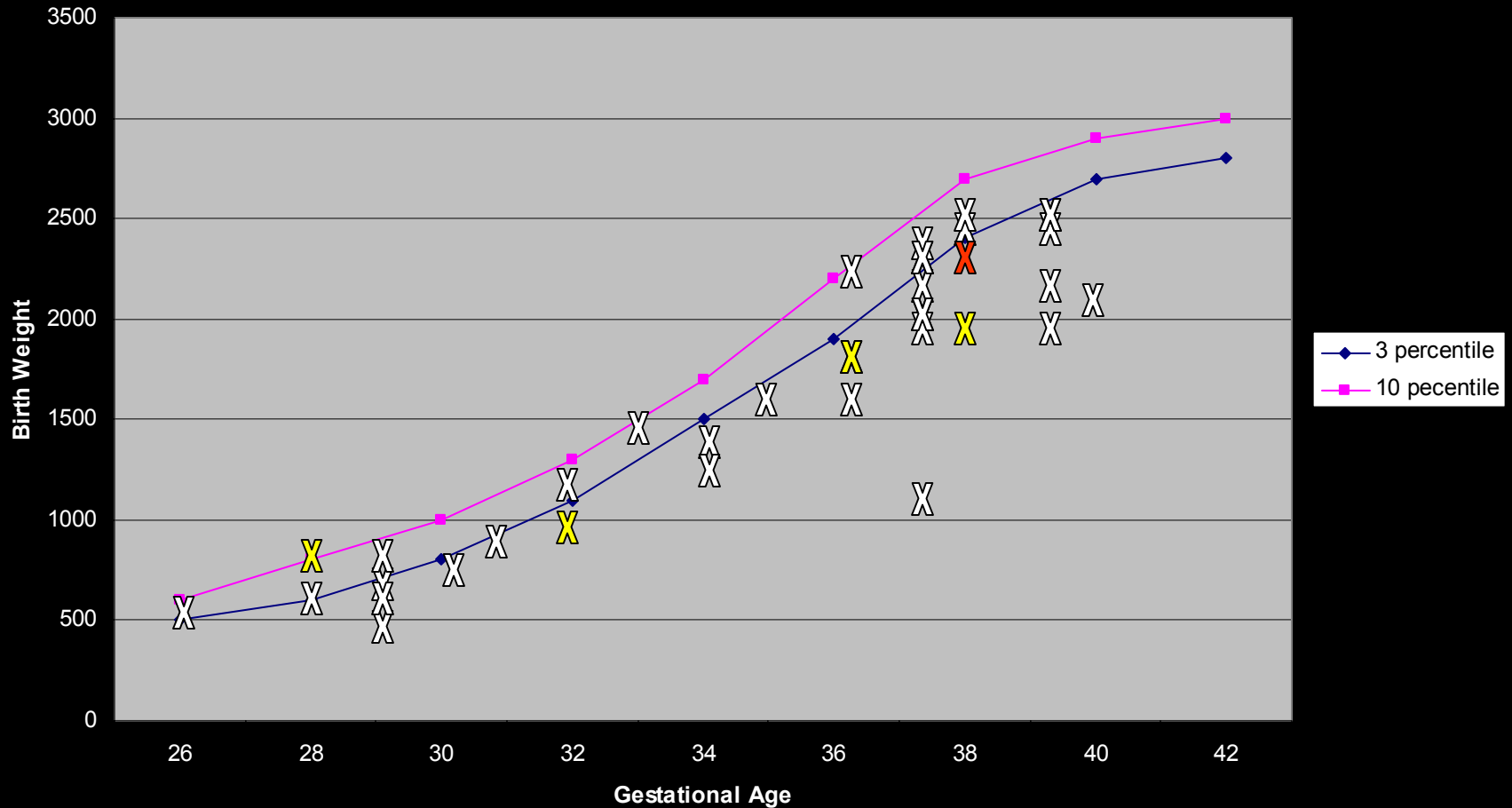
# Hybridization Sites in AGA Placentas



# Birth Weight Distribution - Males

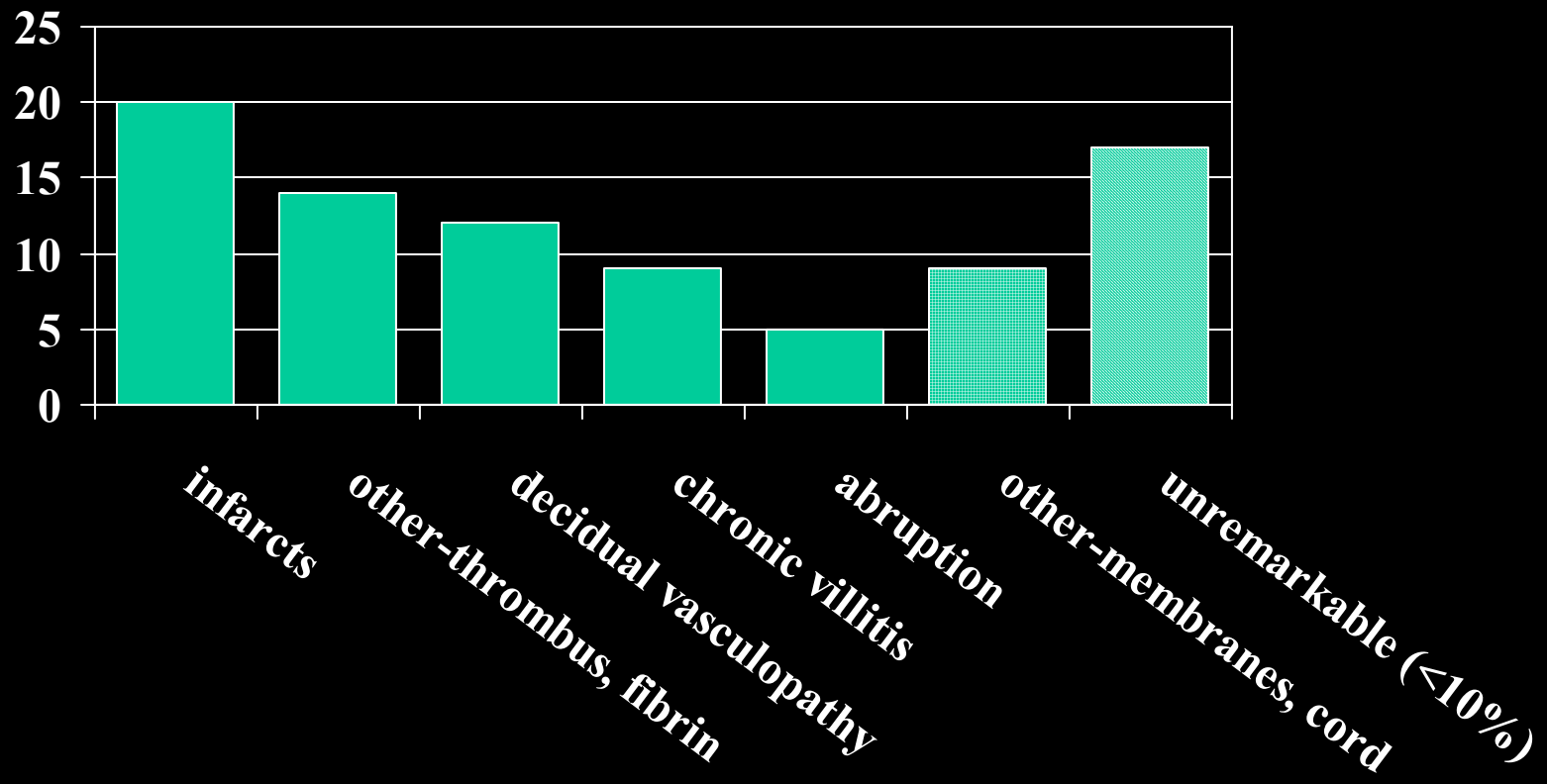


# Birth weight Distribution - Females

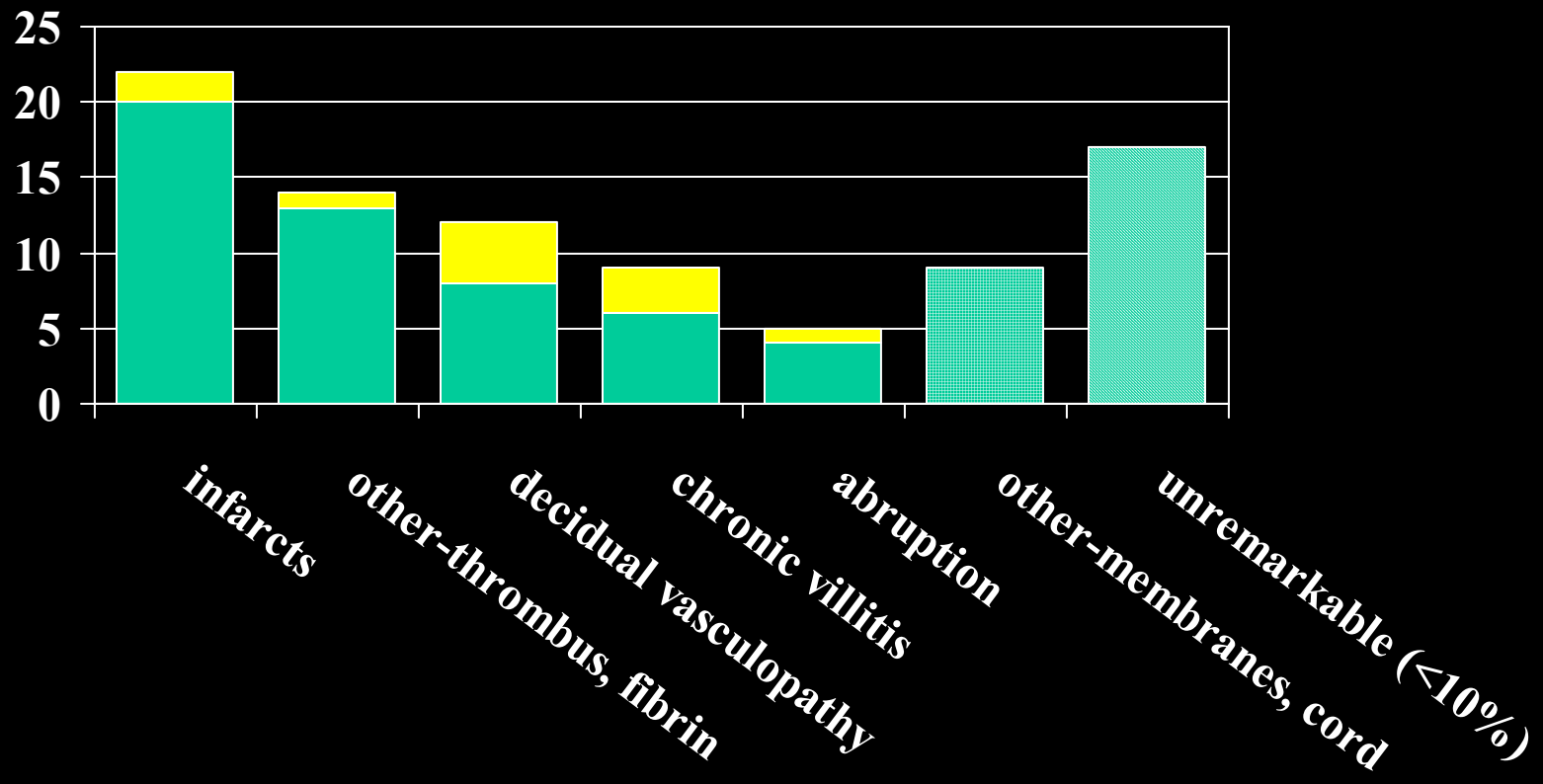




# Placental Histology



# Placental Histology



# Results by molecular testing – UPD among placentas with normal karyotype

- 16 sets of mother / father/ newborn DNA extracted
- All autosomes examined with 1 to 3 dinucleotide repeats
- End point of confirming biparental

# Confirming biparental

- I – infant's polymorphisms only consistent with biparental
- II – consistent with both biparental and uniparental
- III – only consistent with uniparental

# Analyses Performed

- Type I 352 markers
- Type II 704 markers
  - Resolved as biparental on subsequent analyses (additional 1 or 2 markers per chromosome)
- Type III 5 markers

# UPD Results

- Maternal heterodisomy chromosome 14
- Paternal isodisomy chromosome 9
- Nonpaternity

# Case # 235

- Chromosome 14S617

• M	163.1	167.3		
• F	163.1	167.1		
• B	163.1	167.3	M or F	M
biparental or maternal heterodisomy				

- Chromosome 14S587

• M	250.5	261.9		
• F	262.1	265.8		
• B	250.7	261.9	M	M
maternal heterodisomy				

- Chromosome 14S308

• M	201.0	205.1		
• F	204.8	204.8		
• B	201.0	205.0	M	M
maternal heterodisomy				

# Case # 236

- Chromosome 9S930

– M	289.9	289.9		
– F	290.5	298.4		
– B	290.7	290.7	F	F
– paternal isodisomy				

- Chromosome 9S921

– M	174.6	174.6		
– F	196.5	200.60		
– B	200.6	200.6		
– paternal isodisomy			F	F

- Chromosome 9S921

– M	175.0	175.0		
– F	197.0	201.1		
– B	201.1	201.1	F	F
– paternal isodisomy				



# Clinical Outcomes with UPD

- Maternal chromosome 14
  - 38 week infant at 2200 grams
  - Placenta notable for infarcts, villitis
- Paternal chromosome 9
  - 29 week infant at 660 grams
  - Placenta notable for infarcts

# Conclusions

- CPM in 6/75 (8.0 %) well defined IUGR infants versus 1/75 (1.3 %) controls
- No consistent clinical characterization of antepartum complications or placental pathology
- UPD either itself or as a reflection of hidden CPM may play a minor role among infants with IUGR